

## PATENT ABSTRACTS OF JAPAN

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## (54) FLUORESCENT PROTEIN

## (57)Abstract:

PROBLEM TO BE SOLVED: To obtain a fluorescent protein capable of being expressed even by the culture of a host cell at a high temperature (37°C), emitting stronger fluorescent light than those of conventional fluorescent proteins (GFP), and useful as a labeling agent for the analyses of protein localization in live cells, a reporter for the analyses of promoters, etc., by introducing two mutation amino acids into a wild type GFP.

SOLUTION: This fluorescent protein is obtained by mutating the No. 147 serine and the No. 65 serine of the cDNA of a wild type GFP with proline and threonine, respectively, by a site-specific mutation method, etc., transforming Escherichia coil with a plasmid containing the obtained GFPcDNA and subsequently expressing the mutated GFP containing an amino acid sequence of the formula in the Escherichia coil at a high temperature (37°C). The fluorescent protein emits about three-fold fluorescent light that of S65T mutant,

Met-Ser-Lys-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 1 10 20 30 40 50 60 70 80 90 100  
 Val-Ileu-Ser-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 110 120 130 140 150 160 170 180 190 200  
 Val-Ileu-Ser-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 210 220 230 240 250 260 270 280 290 300  
 Val-Ileu-Ser-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 310 320 330 340 350 360 370 380 390 400

Tyr-Ileu-Ser-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 1 10 20 30 40 50 60 70 80 90 100  
 Val-Ileu-Ser-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 110 120 130 140 150 160 170 180 190 200  
 Val-Ileu-Ser-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 210 220 230 240 250 260 270 280 290 300  
 Val-Ileu-Ser-Gly-Phe-His-Glu-Lys-Phe-His-Gly-Tyr-Phe-Pro-Ile-Glu-Tyr-Val  
 310 320 330 340 350 360 370 380 390 400

is contained in a higher concentration than that of the S65T mutant, when expressed in the cell, and emits the fluorescent light under a high temperature (37°C).

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#### LEGAL STATUS

[Date of request for examination]

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[Date of final disposal for application]

[Patent number]

[Date of registration]

[Number of appeal against examiner's decision of rejection]

[Date of requesting appeal against examiner's decision of rejection]

[Date of extinction of right]

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1. This document has been translated by computer. So the translation may not reflect the original precisely.
2. \*\*\*\* shows the word which can not be translated.
3. In the drawings, any words are not translated.

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**CLAIMS**

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[Claim(s)]

[Claim 1] Fluorescence protein which includes an amino acid sequence of a publication in an array number 2.

[Claim 2] Fluorescence protein with which 1 or some amino acid include deletion and an amino acid sequence (however, the 65th place is a threonine and the 147th place is a proline) replaced or added in an array number 2 in an amino acid sequence of a publication.

[Claim 3] DNA which carries out the code of the fluorescence protein according to claim 1 or 2.

[Claim 4] A vector containing DNA according to claim 3.

[Claim 5] A vector according to claim 4 characterized by having arranged DNA according to claim 3 on a \*-ed promotor's lower stream of a river.

[Claim 6] A host cell holding a vector according to claim 4.

[Claim 7] A manufacture method of fluorescence protein including a process which cultivates a host cell according to claim 6, and collects produced protein according to claim 1 or 2.

[Claim 8] A measuring method of the activity of a \*-ed promotor who introduces a vector according to claim 5 into a host cell, and includes a process in which fluorescence emitted from this cell is detected.

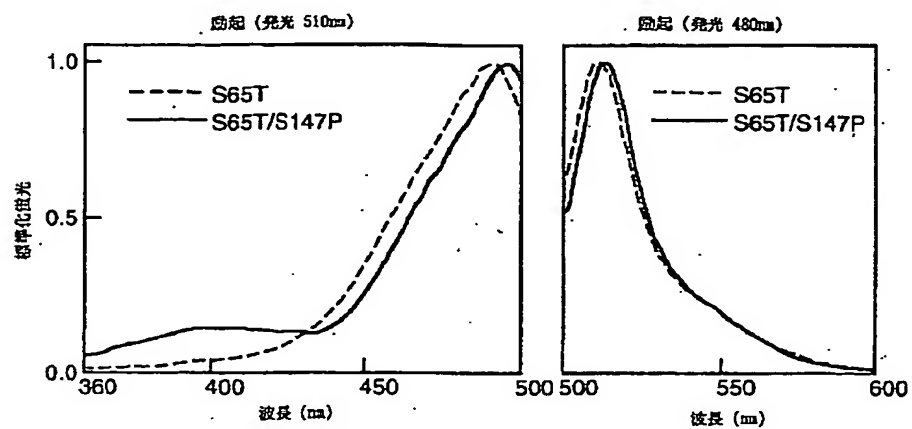
[Claim 9] Fluorescence protein according to claim 1 or 2 characterized by uniting with a \*-ed amino acid sequence.

[Claim 10] How to detect targetting activity in intracellular [ of a \*-ed amino acid sequence ] which introduces fluorescence protein according to claim 9 into a cell, and is characterized by observing distribution in this intracellular one of this fluorescence protein.

[Claim 11] How to detect targetting activity in intracellular [ of a \*-ed amino acid sequence ] which introduces into a host cell a vector in which DNA which carries out the code of the fluorescence protein according to claim 9 was inserted possible [ a manifestation ], and is characterized by observing distribution in this intracellular one of this fluorescence protein.

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Drawing selection **drawing 1**

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ITで消化した「S65T/S147P変異体」のcDNAを挿入し、マウス由来のL cellにカルシウム沈殿法で一過的トランスフェクトした。その細胞を37度で48時間培養した後10%ホルマリンで固定し、蛍光顕微鏡によりノマルスキー(Nomarski)像およびFITCフィルターでの蛍光像(GFPの蛍光)を撮出した(図2A下段)。なお、対照として「S65T変異体」のcDNAを用いた(図2A上段)。この結果、「S65T変異体」と比較して、「S65T/S147P変異体」を発現する細胞は、より明るい蛍光像を示した(図2A右下)。

【0037】また、観察した細胞の内、蛍光を発する細胞の割合、及び細胞の蛍光の強さを測定した(図2B)。図の横軸は、最も蛍光の強かった細胞の蛍光強度を1とした場合における「S65T/S147P」は「S65T/S147P」は蛍光強度を示し、図の縦軸は、蛍光細胞の細胞数を示す。

【0038】この結果、「S65T/S147P変異体」のcDNAを挿入された細胞では、対照と比較して、より高い割合で細胞が蛍光を発していた。また、蛍光強度も対照と比較して顕著に高かった。

【0039】

【発明の効果】本発明により野生型GFPの65番目と147番目のアミノ酸がそれぞれトレオニン、プロリンに置換されたタンパク質が提供された。本発明のタンパク質は、37℃の温度条件下においても蛍光型となり、また従来広く用いられてきた改良型GFPの約3倍の強い蛍光を発 \*

\*すると共に可溶性タンパク質としての発現量も2倍程度増加しているため、従来のタイプに比べ結果として37℃で約5倍程度明るい蛍光を発することが明らかとなった。この改良型GFPは従来のものに比べ37℃での差が顕著であること、微生物のみならず動物細胞でも適用可能であることから、特に動物細胞や幅広い温度で生育可能な酵母などに有効と考えられる。本発明のGFPは、タンパク質の標識として用い、生細胞における分子の局在を観察する目的に適用しているだけでなく、プロモーター解析におけるレポータータンパク質として、またタンパク質の高次構造変化のマーカーとしても有効と考えられ、今後広く細胞生物学、遺伝子工学分野においての利用が期待される。

【0040】

【配列表】

配列番号 : 1

配列の長さ : 717

配列の型 : 核酸

鎖の数 : 二本鎖

20 トポロジー : 直鎖状

配列の塩基 : cDNA to mRNA

配列の特徴

特徴を表す記号 : CDS

存在位置 : 1..714

特徴を決定した方法 : E

配列

ATG AGT AAA GGA GAA CAA CTT TTC ACT CGA GTT GTC CCA ATT CTT GTT	48
Met Ser Lys Gly Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val	
1 5 10 15	
GAA TTA GAT GGT GAT GTT AAT GGG CAC AAA TTT TCT GTC ACT GGA CAG	96
Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu	
20 25 30	
GGT GAA GGT GAT GCA ACA TAC GGA AAA CTT ACC CTT AAA TTT ATT TCC	144
Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys	
35 40 45	
ACT ACT GGA AAA CTA CCT GTT CCA TGG CCA ACA CTT GTC ACT ACT TTC	192
Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe	
50 55 60	
TCT TAT GGT GTT CAA TCC TTT TCA AGA TAC CCA GAT CAT ATG AAA CCG	240
Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg	
65 70 75 80	
CAT GAC TTT TTC AAG AGT GCG ATG CCC GAA GGT TAT GTA CAG GAA AGA	288
His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg	
85 90 95	
ACT ATA TTT TTC AAA GAT GAC GCG AAC TAC AAG ACA GGT GCT GAA GTC	336
Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val	
100 105 110	
AAG TTT GAA GGT GAT ACC CTT GTT AAT AGA ATC GAG TTA AAA GGT ATT	384
Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile	

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11		12	
115	120	125	
GAT TTT AAA GAA GAT CGA AAC ATT CTT CGA CAC AAA TTG GAA TAC AAC			432
Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn			
130	135	140	
TAT AAC TCA CAC AAT GTA TAC ATC ATG CGA CAC AAA CAA AAG AAT CGA			480
Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly			
145	150	155	160
ATC AAA GTT AAC TTC AAA ATT AGA CAC AAC ATT GAA GAT CGA AGC GTT			528
Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val			
165	170	175	
CAA CTA CGA GAC CAT TAT CAA CAA AAT ACT CGA ATT GCG GAT GCG CCT			576
Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro			
180	185	190	
GTC CTT TTA CGA GAC AAC CAT TAC CTG TCC ACA CAA TCT GCC CTT TCG			624
Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser			
195	200	205	
AAA GAT CCC AAC GAA AAG ACA GAC CAC ATG GTC CTT CTT GAG TTT GTA			672
Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val			
210	215	220	
ACA GCT GCT GCG ATT ACA CAT GCG ATG GAT CAA CTA TAC AAA			714
Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys			
225	230	235	
TAA			717

配列番号 : 2

配列の長さ : 238

配列の型 : アミノ酸

\*トポロジー : 直鎖状

配列の種類 : タンパク質

\*

配列

Met Ser Lys Gly Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val
1 5 10 15
Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu
20 25 30
Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys
35 40 45
Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe
50 55 60
Thr Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg
65 70 75 80
His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg
85 90 95
Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val
100 105 110
Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile
115 120 125
Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn
130 135 140
Tyr Asn Pro His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly
145 150 155 160
Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val
165 170 175
Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro

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180 185 190  
Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser  
195 200 205  
Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val  
210 215 220  
Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys  
225 230 235

配列番号 : 3

\* 配列の種類 : cDNA to mRNA

配列の長さ : 717

配列の特徴

配列の型 : 核酸

10 特徴を表す記号 : CDS

鎖の数 : 二本鎖

存在位置 : 1..714

トポロジー : 直鎖状

\* 特徴を決定した方法 : E

配列

ATG AGT AAA GGA GAA GAA CTT TTC ACT GGA GTT GTC CCA ATT CTT GTT	48
Met Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val	
1 5 10 15	
GAA TTA GAT GGT GAT GTT AAT GCG CAC AAA TTT TCT GTC AGT GGA CAG	96
Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu	
20 25 30	
GCT GAA GGT GAT CCA ACA TAC GGA AAA CTT ACC CTT AAA TTT ATT TCC	144
Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys	
35 40 45	
ACT ACT GGA AAA CTA CCT GTT CCA TGG CCA ACA CTT GTC ACT ACT TTC	192
Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe	
50 55 60	
ACT TAT GGT GTT CAA TGC TTT TCA AGA TAC CCA GAT CAT ATG AAA CCG	240
Thr Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg	
65 70 75 80	
CAT GAC TTT TTC AAG AGT GGC ATG CCC GAA GGT TAT GTA CAG GAA AGA	288
His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg	
85 90 95	
ACT ATA TTT TTC AAA CAT GAC GCG AAC TAC AAG ACA CGT GCT GAA GTC	336
Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val	
100 105 110	
AAG TTT GAA GGT GAT ACC CTT GTT AAT AGA ATC GAG TTA AAA GGT ATT	384
Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile	
115 120 125	
GAT TTT AAA GAA GAT GGA AAC ATT CTT GGA CAC AAA TTG GAA TAC AAC	432
Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn	
130 135 140	
TAT AAC CCA CAC AAT GTA TAC ATC ATG CCA CAC AAA CAA AAG AAT GGA	480
Tyr Asn Pro His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly	
145 150 155 160	
ATC AAA GTT AAC TTC AAA ATT AGA CAC AAC ATT GAA CAT GGA AGC GTT	528
Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val	
165 170 175	
CAA CTA GCA GAC CAT TAT CAA CAA AAT ACT GCA ATT GGC GAT GGC CCT	576
Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro	
180 185 190	
GTC CTT TTA CCA GAC AAC CAT TAC CTG TCC ACA CAA TCT GGC CTT TCG	624

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符号平 1 0 - 2 3 4 3 8 2

15 16  
Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser  
195 205 205  
AAA GAT CCC AAC GAA AAG AGA GAC CAC ATG GTC CTT CTT GAG TTT GTA 672  
Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val  
210 215 220  
ACA GCT GCT GCG ATT ACA CAT GCG ATG GAT GAA CTA TAC AAA 714  
Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys  
225 230 235  
TAA 717

配列番号 : 4

配列の長さ : 36

配列の型 : 核酸

10\*鎖の数 : 一本鎖

トポロジー : 直鎖状

\* 配列の種類 : 他の核酸 合成DNA

配列

GGCCCCGAT CCATGAGTAA AGGAGAAGAA CTTTTC 36

配列番号 : 5

配列の長さ : 39

配列の型 : 核酸

※鎖の数 : 一本鎖

トポロジー : 直鎖状

※ 配列の種類 : 他の核酸 合成DNA

配列

GGCCAGGTA CCTTATTTGT ATAGTTCATC CATGCCATG 39

配列番号 : 6

配列の長さ : 31

配列の型 : 核酸

29★鎖の数 : 一本鎖

トポロジー : 直鎖状

★ 配列の種類 : 他の核酸 合成DNA

配列

TTCACCCGGG ATGAGTAAAG GAGAAGAACT T 31

配列番号 : 7

配列の長さ : 33

配列の型 : 核酸

☆鎖の数 : 一本鎖

トポロジー : 直鎖状

☆ 配列の種類 : 他の核酸 合成DNA

配列

GCACGAATTC TATTGTATA GTTCATCCAT GCC 33

【図面の簡単な説明】

【図1】「S65T/S147P変異体」及び「S65T変異体」の励起・蛍光スペクトルの測定結果を示す図である。

【図2】図2Aは、「S65T/S147P変異体」及び「S65T変異体」のcDNAが導入された細胞を蛍光顕微鏡により検出◆

30◆し、そのノマルスキー像及び蛍光像を示す顕微鏡写真である。図2Bは、被検細胞の中で蛍光を発する細胞の割合及びその細胞の蛍光の強さの測定結果を示す図である。

【図1】

